

RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical
Information Center (STIC) no errors detected.

Application Serial Number: 09/930, 440c
Source: EFW/6
Date Processed by STIC: 12-14-04

ENTERED



IFW16

RAW SEQUENCE LISTING

DATE: 12/14/2004

PATENT APPLICATION: US/09/930,440C

TIME: 14:00:39

Input Set : A:\03940077pa.txt

Output Set: N:\CRF4\12142004\I930440C.raw

3 <110> APPLICANT: Betenbaugh, Michael J.
 4 Lawrence, Shawn J.
 5 Lee, Yuan C.
 6 Coleman, Timothy A.
 8 <120> TITLE OF INVENTION: Engineering Intracellular Sialylation Pathways
 10 <130> FILE REFERENCE: 03940077pa
 12 <140> CURRENT APPLICATION NUMBER: 09/930,440C
 13 <141> CURRENT FILING DATE: 2001-08-16
 15 <150> PRIOR APPLICATION NUMBER: US 60/122,582
 16 <151> PRIOR FILING DATE: 1999-03-02
 18 <150> PRIOR APPLICATION NUMBER: US 60/169,624
 19 <151> PRIOR FILING DATE: 1999-12-08
 21 <150> PRIOR APPLICATION NUMBER: US 60/227,579
 22 <151> PRIOR FILING DATE: 2000-08-25
 24 <150> PRIOR APPLICATION NUMBER: US 09/516,793
 25 <151> PRIOR FILING DATE: 2000-03-01
 27 <160> NUMBER OF SEQ ID NOS: 18
 29 <170> SOFTWARE: PatentIn version 3.2
 31 <210> SEQ ID NO: 1
 32 <211> LENGTH: 1429
 33 <212> TYPE: DNA
 34 <213> ORGANISM: Homo sapiens
 36 <400> SEQUENCE: 1

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39 gagaatggag	aaatcaactt	ttcagtaatt	ggtcagtatg	tggattatct	tgtgaaagaa	120
41 caggagtgta	agaacatttt	tgtgaatggc	acaacaggag	aaggcctgtc	cctgagcgtc	180
43 tcagagcgtc	gccagggtgc	agaggagtgg	gtgacaaaag	ggaaggacaa	gctggatcag	240
45 gtgataattc	acgtaggagc	actgagcttg	aaggagtcac	aggaactggc	ccaacatgca	300
47 gcagaaatag	gagctgatgg	catcgctgtc	attgcaccgt	tcttcctcaa	gccatggacc	360
49 aaagatatcc	tgattaattt	cctaaaggaa	gtggctgctg	ccgcccctgc	cctgccattt	420
51 tattactatc	acattcctgc	cttgacaggg	gtaaagattc	gtgctgagga	gttggtggat	480
53 gggattctgg	ataagatccc	caccttccaa	gggctgaaat	tcagtgatac	agatctctta	540
55 gacttcgggc	aatgtgttga	tcagaatcgc	cagcaacagt	ttgctttcct	ttttggggtg	600
57 gatgagcaac	tgttgagtgc	tctggtgatg	ggagcaactg	gagcagtggg	cagttttgta	660
59 tccagagatt	tatcaacttt	gttgtcaaac	taggttttgg	agtgtcacag	accaaagcca	720
61 tcatgactct	ggtctctggg	attccaatgg	gccaccccg	gcttccactg	cagaaagcct	780
63 ccagggagtt	tactgatagt	gctgaagcta	aactgaagag	cctggatttc	ctttctttca	840
65 ctgattttaa	ggatggaaac	ttggaagctg	gtagctagtg	cctctctatc	aaatcagggt	900
67 ttgcaccttg	agacataatc	taccttaaat	agtgcatttt	tttctcaggg	aattttgat	960
69 gaacttgaat	aaactctcct	agcaaatgaa	atctcacaat	aagcattgag	gtaccttttg	1020
71 tgagccttaa	aaagtcttat	tttgtgaagg	ggcaaaaact	ctaggagtca	caactctcag	1080
73 tcattcattt	cacagatttt	tttgtggaga	aatttctgtt	tatatggatg	aaatggaatc	1140
75 aagaggaaaa	ttgtaattga	ttaattccat	ctgtcttttag	gagctctcat	tatctcggtc	1200

(pg. 6)

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77 tctgggttcoct aatcctatatt taaagttgtc taatttttaa ccactataat atgtcttcat 1260
79 ttttaataaat attcatttgg aatctaggaa aactctgagc tactgcattt aggcaggcac 1320
81 ttttaataacca aactgtaaca tgtctcaact gtatacaact caaaatacac cagctcattt 1380
83 ggctgctcag tctaactcta gaatggatgc ttttgaattc atttcgatg 1429
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87 <211> LENGTH: 304
88 <212> TYPE: PRT
89 <213> ORGANISM: Homo sapiens
91 <400> SEQUENCE: 2
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97 Thr Pro Met Thr Glu Asn Gly Glu Ile Asn Phe Ser Val Ile Gly Gln
98 20 25 30
101 Tyr Val Asp Tyr Leu Val Lys Glu Gln Gly Val Lys Asn Ile Phe Val
102 35 40 45
105 Asn Gly Thr Thr Gly Glu Gly Leu Ser Leu Ser Val Ser Glu Arg Arg
106 50 55 60
109 Gln Val Ala Glu Glu Trp Val Thr Lys Gly Lys Asp Lys Leu Asp Gln
110 65 70 75 80
113 Val Ile Ile His Val Gly Ala Leu Ser Leu Lys Glu Ser Gln Glu Leu
114 85 90 95
117 Ala Gln His Ala Ala Glu Ile Gly Ala Asp Gly Ile Ala Val Ile Ala
118 100 105 110
121 Pro Phe Phe Leu Lys Pro Trp Thr Lys Asp Ile Leu Ile Asn Phe Leu
122 115 120 125
125 Lys Glu Val Ala Ala Ala Ala Pro Ala Leu Pro Phe Tyr Tyr Tyr His
126 130 135 140
129 Ile Pro Ala Leu Thr Gly Val Lys Ile Arg Ala Glu Glu Leu Leu Asp
130 145 150 155 160
133 Gly Ile Leu Asp Lys Ile Pro Thr Phe Gln Gly Leu Lys Phe Ser Asp
134 165 170 175
137 Thr Asp Leu Leu Asp Phe Gly Gln Cys Val Asp Gln Asn Arg Gln Gln
138 180 185 190
141 Gln Phe Ala Phe Leu Phe Gly Val Asp Glu Gln Leu Leu Ser Ala Leu
142 195 200 205
145 Val Met Gly Ala Thr Gly Ala Val Gly Ser Phe Val Ser Arg Asp Leu
146 210 215 220
149 Ser Thr Leu Leu Ser Asn Val Leu Glu Cys His Arg Pro Lys Pro Ser
150 225 230 235 240
153 Leu Trp Ser Leu Gly Phe Gln Trp Ala His Pro Gly Phe His Cys Arg
154 245 250 255
157 Lys Pro Pro Gly Ser Leu Leu Ile Val Leu Lys Leu Asn Arg Ala Trp
158 260 265 270
161 Ile Ser Phe Leu Ser Leu Ile Arg Met Glu Thr Trp Lys Leu Val Ala
162 275 280 285
165 Ser Ala Ser Leu Ser Asn Gln Gly Phe Ala Pro Leu Arg His Asn Leu
166 290 295 300
169 <210> SEQ ID NO: 3
170 <211> LENGTH: 1305

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Input Set : A:\03940077pa.txt

Output Set: N:\CRF4\12142004\I930440C.raw

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171 <212> TYPE: DNA
172 <213> ORGANISM: Homo sapiens
174 <400> SEQUENCE: 3
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177 cggggcgccg cgcggaagct gcagcgcaac tctcgcgccg gccagggccg aggtgtggag      120
179 aagccccgcg acctggcagc cctaattctg gcccggggag gcagcaaagg catccccctg      180
181 aagaacatta agcacctggc ggggggtccc ctcattggct gggctcctgcg tgcggccctg      240
183 gattcagggg ccttcagag tgtatgggtt tcgacagacc atgatgaaat tgagaatgtg      300
185 gccaaacaat ttggtgcaca agttcatcga agaagttctg aagtttcaaa agacagctct      360
187 acctcactag atgccatcat agaatttctt aattatyata atgaggktga cattgtagga      420
189 aatattcaag ctacttctyc atgtttacat cctactgata ttcaaaaagt tgcagaaatg      480
191 attcgagaag aaggatatga ttctgktttc tctgttgtga gacgccatca gtttcgatgg      540
193 agtgaaattc agaaaggagt tcgtgaagtg accgaacctc tgaatttaaa tccagctaaa      600
195 cggcctcgtc gacaagactg ggatggagaa ttatatgaaa atggctcatt ttattttgct      660
197 aaaagacatt tgatagagat gggttacttg cagggtggaa aaatggcata ctacgaaatg      720
199 cgagctgaac atagtgtgga tatagatgtg gatattgatt ggcctattgc agagcaaaga      780
201 gtattaagat atggctatth tggcaaagag aagcttaagg aaataaaaact tttggtttgc      840
203 aatattgatg gatgtctcac caatggccac atttatgtat caggagacca aaaagaaata      900
205 atatcttatg atgtaaaaga tgctattggg ataagtttat taaagaaaag tggatttgag      960
207 gtgaggctaa tctcagaaa ggctgtttca aagcagacgc tgtcttcttt aaaactggat      1020
209 tgcaaaatgg aagtcagtgt atcagacaag ctagcagttg tagatgaatg gagaaaagaa      1080
211 atgggcctgt gctggaaaag agtggcatat cttggaaatg aagtgtctga tgaagagtgc      1140
213 ttgaagagag tgggcctaag tggcgctcct gctgatgcct gttcctacgc ccagaaggct      1200
215 gttggataca ttgcaaagt taatgggtggc cgtgggtgcc tccgagaatt tgcagagcac      1260
217 atttgcctac taatggaaaa agttaataat tcatgccaaa aatag      1305
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221 <211> LENGTH: 434
222 <212> TYPE: PRT
223 <213> ORGANISM: Homo sapiens
226 <220> FEATURE:
227 <221> NAME/KEY: misc_feature
228 <222> LOCATION: (133)..(133)
229 <223> OTHER INFORMATION: Xaa can be any naturally occurring amino acid
231 <220> FEATURE:
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233 <222> LOCATION: (136)..(136)
234 <223> OTHER INFORMATION: Xaa can be any naturally occurring amino acid
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238 <222> LOCATION: (147)..(147)
239 <223> OTHER INFORMATION: Xaa can be any naturally occurring amino acid
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243 <222> LOCATION: (169)..(169)
244 <223> OTHER INFORMATION: Xaa can be any naturally occurring amino acid
246 <400> SEQUENCE: 4
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252 Gly Arg Pro Ser Arg Gly Arg Pro Pro Lys Leu Gln Arg Asn Ser Arg

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253          20          25          30
256 Gly Gly Gln Gly Arg Gly Val Glu Lys Pro Pro His Leu Ala Ala Leu
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260 Ile Leu Ala Arg Gly Gly Ser Lys Gly Ile Pro Leu Lys Asn Ile Lys
261          50          55          60
264 His Leu Ala Gly Val Pro Leu Ile Gly Trp Val Leu Arg Ala Ala Leu
265 65          70          75          80
268 Asp Ser Gly Ala Phe Gln Ser Val Trp Val Ser Thr Asp His Asp Glu
269          85          90          95
272 Ile Glu Asn Val Ala Lys Gln Phe Gly Ala Gln Val His Arg Arg Ser
273          100          105          110
276 Ser Glu Val Ser Lys Asp Ser Ser Thr Ser Leu Asp Ala Ile Ile Glu
277          115          120          125
W--> 280 Phe Leu Asn Tyr Xaa Asn Glu Xaa Asp Ile Val Gly Asn Ile Gln Ala
281          130          135          140
W--> 284 Thr Ser Xaa Cys Leu His Pro Thr Asp Leu Gln Lys Val Ala Glu Met
285 145          150          155          160
W--> 288 Ile Arg Glu Glu Gly Tyr Asp Ser Xaa Phe Ser Val Val Arg Arg His
289          165          170          175
292 Gln Phe Arg Trp Ser Glu Ile Gln Lys Gly Val Arg Glu Val Thr Glu
293          180          185          190
296 Pro Leu Asn Leu Asn Pro Ala Lys Arg Pro Arg Arg Gln Asp Trp Asp
297          195          200          205
300 Gly Glu Leu Tyr Glu Asn Gly Ser Phe Tyr Phe Ala Lys Arg His Leu
301          210          215          220
304 Ile Glu Met Gly Tyr Leu Gln Gly Gly Lys Met Ala Tyr Tyr Glu Met
305 225          230          235          240
308 Arg Ala Glu His Ser Val Asp Ile Asp Val Asp Ile Asp Trp Pro Ile
309          245          250          255
312 Ala Glu Gln Arg Val Leu Arg Tyr Gly Tyr Phe Gly Lys Glu Lys Leu
313          260          265          270
316 Lys Glu Ile Lys Leu Leu Val Cys Asn Ile Asp Gly Cys Leu Thr Asn
317          275          280          285
320 Gly His Ile Tyr Val Ser Gly Asp Gln Lys Glu Ile Ile Ser Tyr Asp
321          290          295          300
324 Val Lys Asp Ala Ile Gly Ile Ser Leu Leu Lys Lys Ser Gly Ile Glu
325 305          310          315          320
328 Val Arg Leu Ile Ser Glu Arg Ala Cys Ser Lys Gln Thr Leu Ser Ser
329          325          330          335
332 Leu Lys Leu Asp Cys Lys Met Glu Val Ser Val Ser Asp Lys Leu Ala
333          340          345          350
336 Val Val Asp Glu Trp Arg Lys Glu Met Gly Leu Cys Trp Lys Glu Val
337          355          360          365
340 Ala Tyr Leu Gly Asn Glu Val Ser Asp Glu Glu Cys Leu Lys Arg Val
341          370          375          380
344 Gly Leu Ser Gly Ala Pro Ala Asp Ala Cys Ser Tyr Ala Gln Lys Ala
345 385          390          395          400
348 Val Gly Tyr Ile Cys Lys Cys Asn Gly Gly Arg Gly Ala Ile Arg Glu
349          405          410          415

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361 <211> LENGTH: 1080
362 <212> TYPE: DNA
363 <213> ORGANISM: Homo sapiens
365 <400> SEQUENCE: 5
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370 cgcattggcca aggagtgtgg ggctgattgt gccagttcc agaagagtga gctagaattc      180
372 aagtttaatc ggaaagcctt ggagaggcca tacacctga agcattcctg ggggaagacg      240
374 tacggggagc acaaacgaca tctggagttc agccatgacc agtacaggga gctgcagagg      300
376 tacgccgagg aggttgggat cttcttcact gcctctggca tggatgagat ggcagttgaa      360
378 ttcttgcata aactgaatgt tccatttttc aaagttggat ctggagacac taataatttt      420
380 ctttatctgg aaaagacagc caaaaaaggt cgcccaatgg tgatctccag tgggatgcag      480
382 tcaatggaca ccatgaagca agtttatcag atcgtgaagc ccctcaacc caacttctgc      540
384 ttcttgcagt gtaccagcgc ataccgcctc cagcctgagg acgtcaacct gcgggtcatc      600
386 tcggaatata agaagctctt tcttgacatt cccatagggt attctgggca tgaaacaggc      660
388 atagcgatat ctgtggccgc agtggctctg ggggccaaag tgttggaaac tcacataact      720
390 ttggacaaga cctggaaggg gagtgaccac tcggcctcgc tggagcctgg agaactggcc      780
392 gagctgggtg ggtcagtgcg tcttgtggag cgtgccctgg gctccccaac caagcagctg      840
394 ctgccctgtg agatggcctg caatgagaag ctgggcaagt ctgtggtggc caaagtgaag      900
396 attccggaag gcaccattct aacaatggac atgctcaccg tgaaggtggg tgagcccaaa      960
398 gcctatcctc ctgaagacat ctttaattcta gtgggcaaga aggtcctggt cactgttgaa     1020
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404 <211> LENGTH: 359
405 <212> TYPE: PRT
406 <213> ORGANISM: Homo sapiens
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414 His Pro Cys Phe Ile Ile Ala Glu Ile Gly Gln Asn His Gln Gly Asp
415           20           25           30
418 Leu Asp Val Ala Lys Arg Met Ile Arg Met Ala Lys Glu Cys Gly Ala
419           35           40           45
422 Asp Cys Ala Lys Phe Gln Lys Ser Glu Leu Glu Phe Lys Phe Asn Arg
423           50           55           60
426 Lys Ala Leu Glu Arg Pro Tyr Thr Ser Lys His Ser Trp Gly Lys Thr
427 65           70           75           80
430 Tyr Gly Glu His Lys Arg His Leu Glu Phe Ser His Asp Gln Tyr Arg
431           85           90           95
434 Glu Leu Gln Arg Tyr Ala Glu Glu Val Gly Ile Phe Phe Thr Ala Ser
435           100          105          110
438 Gly Met Asp Glu Met Ala Val Glu Phe Leu His Glu Leu Asn Val Pro
439           115          120          125
442 Phe Phe Lys Val Gly Ser Gly Asp Thr Asn Asn Phe Pro Tyr Leu Glu
443           130          135          140

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RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/09/930,440C

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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:4; Xaa Pos. 133,136,147,169
Seq#:9; N Pos. 1,3,6,12,13,15,18
Seq#:10; N Pos. 3,4,6,9,10,11,12,15,17,18
Seq#:11; N Pos. 3,4,6,9,10,11,12,15,17,18
Seq#:12; N Pos. 2,3,6,9,12,15,16,17,18

Invalid <213> Response:

Use of "Artificial" only as "<213> Organism" response is incomplete,
per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:9,10,11,12,13,14,15,16

VERIFICATION SUMMARY

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L:280 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:128
L:284 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:144
L:288 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:160
L:686 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:0
L:750 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:0
L:814 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:0
L:873 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12 after pos.:0